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Sara Murillo-Sánchez, Damien Beaufils, Juan Manuel González Mañas, Robert Pascal, Kepa Ruiz-Mirazo, et al.. Fatty acids' double role in the prebiotic formation of a hydrophobic dipeptide. *Chemical Science*, 2016, 7 (5), pp.3406-3413. 10.1039/c5sc04796j . hal-03549304

HAL Id: hal-03549304

<https://hal.science/hal-03549304>

Submitted on 31 Jan 2022

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Cite this: *Chem. Sci.*, 2016, 7, 3406

Fatty acids' double role in the prebiotic formation of a hydrophobic dipeptide†

Sara Murillo-Sánchez,^a Damien Beaufile,^b Juan Manuel González Mañas,^c Robert Pascal^{*b} and Kepa Ruiz-Mirazo^{*ad}

In search of a connection between prebiotic peptide chemistry and lipid compartments, the reaction of a 5(4*H*)-oxazolone with leucinamide was extensively explored under buffered aqueous conditions, where diverse amphiphiles and surfactants could form supramolecular assemblies. Significant increases in yield and changes in stereoselectivity were observed when fatty acids exceeded their critical aggregation concentration, self-assembling into vesicles in particular. This effect does not take place below the fatty acid solubility limit, or when other anionic amphiphiles/surfactants are used. Data from fluorimetric and Langmuir trough assays, complementary to the main HPLC results reported here, demonstrate that the dipeptide product co-localizes with fatty acid bilayers and monolayers. Additional experiments in organic solvents suggest that acid–base catalysis operates at the water–aggregate interface, linked to the continuous proton exchange dynamics that fatty acids undergo at pH values around their effective pK_a . These simple amphiphiles could therefore play a dual role as enhancers of peptide chemistry under prebiotic conditions, providing soft and hydrophobic organic domains through self-assembly and actively inducing catalysis at their interface with the aqueous environment. Our results support a systems chemistry approach to life's origin.

Received 11th December 2015
Accepted 9th February 2016

DOI: 10.1039/c5sc04796j

www.rsc.org/chemicalscience

Introduction

Chemistry today is up to the challenge of working with increasingly diverse combinations of molecules and supramolecular structures, both under quasi-equilibrium and non-equilibrium conditions, providing new means to explain their complex dynamic behaviour. The field of 'systems chemistry' has been launched precisely to address this challenge^{1,2} with strong implications for studies in connection with biology and, in particular, with research on the origins of life.³ In this context, the idea that the chemical roots of biological phenomena cannot be found in the properties of a single type of molecule (often assumed to be a biopolymer, like RNA), but must be approached in a more encompassing way that includes reaction networks made of different bio-molecular precursors, is gaining momentum. This novel view has recently received

further support from evidence that there could be a common prebiotic chemistry underlying the synthesis of a rich variety of these precursors.⁴ The obvious next step would be to determine the way in which some of these initial biomonomers engage in interactions that contribute to producing and maintaining higher levels of molecular complexity and diversity, as we actually observe in biology. Here we report experimental results demonstrating that a primitive synergy of this kind could occur between two such types of molecules, namely those whose prebiotic relevance is better grounded: amino acids, as peptide and protein precursors,^{5,6} and fatty acids, as lipid precursors.^{7,8}

Previous studies have shown that the activation of amino acid monomers or peptides can lead to their oligomerization in aqueous solution.^{9,10} Peptide oligomers are formed by the stepwise polymerization of α -amino acid *N*-carboxyanhydrides (NCAs), as prebiotically plausible activated monomers. In addition to this pathway, the activation of the C-terminus in peptides or acylated amino acids has been shown to proceed through the formation of 5(4*H*)-oxazolones from reagents compatible with early Earth conditions.¹¹ In turn, fatty acids, as single-chain amphiphilic precursors of the standard (but remarkably more complex) phospholipids found in bacterial and eukaryote biomembranes, were long ago shown to self-assemble into vesicles at intermediate pH values.^{12,13} This aggregation state is favoured due to the fact that both the neutral and anionic forms of the fatty acid are present in similar proportions (at $pH \sim pK_a$) and tend to establish H-bonds

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† Electronic supplementary information (ESI) available: Exhaustive data on the outcome of reactions, the aggregation state of the surfactants and the affinity of the dipeptide product for the aggregates. See DOI: 10.1039/c5sc04796j



between them, leading to molecular pairs with cylindrical geometry (see Fig. S1, in the ESI†), which spontaneously assemble into flat bilayers. In recent years, an increasing number of research groups have explored the properties of these supramolecular structures, given their potential as pro-cellular model systems.^{14–22}

However, relatively little effort has been devoted to putting together these two prebiotic systems. Although the idea of combining peptide chemistry with lipid aggregates is not new, it has been implemented mostly in making use of standard liposomes (*i.e.*, vesicles made of phospholipids). For instance, mixing peptides with lipid aggregates was advocated to improve the stability and functionality of the latter.²³ In addition, the polymerization of aminoacyl adenylates into peptides was claimed to yield increased chain lengths in the presence of vesicles,²⁴ even though the mechanism does not correspond to a direct process, but actually involves a prior conversion into NCAs.^{25,26} Luisi's group also performed a series of investigations demonstrating the role of phospholipid compartments in the efficiency and stereoselectivity of peptide polymerization.^{27–30} In an interesting step forward, liposomes made of cationic surfactants were later shown to favour the oligomerization of encapsulated anionic amino acid monomers.³¹ And, more recently, the formation of tryptophan based dipeptides in the aqueous cavity of phosphatidylcholine vesicles has been reported.³²

Much less is found in the literature regarding the use of fatty acid vesicles to perform similar experiments. As an exception, Matsuno's lab performed some experiments in which they reported the oligomerization of encapsulated glycine, not only in phospholipids but also in fatty acid vesicle compartments.³³ More directly relevant to our approach, a couple of years ago, Szostak's group managed to carry out, within oleic acid vesicles and with the catalytic assistance of a pre-encapsulated dipeptide (Ser–His), the synthesis of a strongly hydrophobic peptide, which spontaneously joined the membrane and, thereby, contributed to its growth.³⁴ In this article we point towards a similar idea but approach it from a complementary angle. We show how a prototypical prebiotic synthesis of a hydrophobic dipeptide (starting from precursors expected to be protected by lipophilic environments)³⁵ is significantly enhanced, without the aid of a soluble catalyst, just in the presence of vesicles made of fatty acids. This effect, which relies on the specific fatty acid nature of the compartments, is also partially detected when reactants are put in organic solvents (indicative of direct chemical catalysis by the fatty acids). Thus, our results give fatty acids wider significance as prebiotic compounds than has been commonly assumed in the field of the origins of life.

Materials and methods

Chemicals and general information

N-Ethyl-*N'*-(3-dimethylaminopropyl)-carbodiimide·HCl (EDC), *H*-*L*-Leu-NH₂, *Ac*-*L*-Tyr(Me)-OH and *Boc*-*L*-Tyr(Me)-OH were purchased from Bachem. Piperazine-*N,N'*-bis(3-propanesulfonic acid) (PIPPS) was purchased from CalbioChem. 1,2-Dioleoyl-*sn*-

glycero-phosphate (monosodium salt) was purchased from Avanti Polar Lipids. The rest of the reagents and solvents were purchased from Sigma Aldrich. All compounds were used without further purification. In all experiments pH was monitored using a Thermo Orion 3-STAR pH-meter with a VWR electrode. NMR spectra in DMSO-*d*₆ solution were recorded on a Bruker Avance 300 spectrometer at 300 MHz for ¹H. HPLC analysis (Method A) was carried out on a Waters Alliance system including a E2695 separation module and a 2998 photodiode array detector; column BDS Hypersil C18 (3 μm, 2.1 × 50 mm). The elution method used was: flow 0.2 mL min⁻¹, solvent A 0.1% TFA in H₂O, solvent B 0.1% TFA in CH₃CN, gradient 90% B to 70% over 25 min; detection 273 nm. HPLC-ESI-MS analysis was carried out (Method B) using a Waters Synapt G2-S system connected to Waters Acquity UPLC H-Class apparatus equipped with an Acquity UPLC BEH C18, 1.7 μm 2.1 × 50 mm column; the gradient of solvent A was 0.01% formic acid in H₂O and solvent B: 0.01% formic acid in acetonitrile; flow rate: 0.5 mL min⁻¹; linear gradient 0–100% B over 3 min.

Experimental procedures

4-Methoxybenzyl-2-methyl-5(4*H*)-oxazolone, 2a. The oxazolone was prepared in a racemic form by following a published protocol.³⁶ *Ac*-*L*-Tyr(Me)-OH (500 mg) was dissolved in CH₂Cl₂ (20 mL) under magnetic stirring, at 0 °C. When everything was dissolved, 445 mg of EDC was added and the mixture was stirred for 1 h at 0 °C. The reaction mixture was then diluted with 30 mL of CH₂Cl₂, and washed successively with water (30 mL), saturated aqueous NaHCO₃ (30 mL, twice), and saturated NaCl (30 mL, twice), then dried with Na₂SO₄ and concentrated under reduced pressure. The solid product crystallized upon storage at –20 °C for 24 hours.

4-Isobutyl-2-methyl-5(4*H*)-oxazolone, 2b. The oxazolone was prepared from *Ac*-*D,L*-Leu-OH using a similar protocol³⁷ and exhibited the 2.0 Hz long range coupling constant already described in the literature between the 2-methyl group and the α -proton.³⁸ ¹H NMR (300 MHz, DMSO-*d*₆), δ _H ppm: 4.34 (m, 1H); 2.15 (d, ⁵*J* = 2.0 Hz, 3H); 1.84 (m, 1H); 1.40–1.58 (m, 2H); 0.93 (d, *J* = 6.8 Hz, 3H), 0.91 (d, *J* = 6.8 Hz, 3H).

***Ac*-Tyr(Me)-Leu-NH₂ dipeptides.** The dipeptide diastereomers were prepared from either *Boc*-*D*-Tyr(Me)-OH or *Boc*-*L*-Tyr(Me)-OH. A mixture of the selected *Boc*-protected derivative, hydroxybenzotriazole hydrate (HOBt, 1 eq.) and CH₂Cl₂ (10 mL mmol⁻¹) was stirred at 0 °C for 10 min. Then 1,3-dicyclohexylcarbodiimide (DCC, 1 eq.) was introduced and stirring was continued for 15 min at 0 °C. The solid urea by-product was removed by filtration, twice washed with CH₂Cl₂ and the combined filtrates were concentrated under vacuum without heating. The oily residue was diluted in DMF (5 mL mmol⁻¹), *H*-*L*-Leu-NH₂ hydrochloride (1 eq.) was added with *N,N*-diisopropylethylamine (DIEA, 2 eq.) and the mixture was stirred overnight at r.t. The dipeptide product was precipitated upon addition of 1 M Na₂CO₃ solution and collected by filtration, washed with water and dried under vacuum. The *Boc* protecting group was removed using 90 : 10 v/v TFA/water reagent at r.t. for 30 min. The mixture was concentrated under reduced pressure



and the product was precipitated by the addition of cold ethyl ether, filtered off, washed with ethyl ether and dried under vacuum. Acetylation was performed with acetic anhydride (1.5 eq.) and DIEA (2 eq.) in CH_2Cl_2 for 5 h at 0 °C. The mixture was diluted with water and extracted with ethyl acetate. The organic layers were washed with saturated Na_2CO_3 solution, then with 1 M KHSO_4 and saturated NaCl and finally dried over anhydrous Na_2SO_4 . The solvent was removed under reduced pressure to yield the product as a white solid.

Ac-L-Tyr(Me)-L-Leu-NH₂. ¹H NMR (300 MHz, DMSO-d₆), δ ppm: 8.05 (d, 1H, $J = 8.1$ Hz), 7.92 (d, 1H, $J = 8.4$ Hz), 7.15–7.18 (m, 3H), 6.99 (s, 1H), 6.82 (d, 2H, $J = 4.8$ Hz), 4.43 (m, 1H), 4.20 (q, 1H, $J = 7.5$ Hz), 3.71 (s, 3H), 2.93 (dd, 1H, $J = 13.8$ Hz, $J = 4.5$ Hz), 2.66 (dd, 1H, $J = 9.9$ Hz, $J = 13.8$ Hz), 1.76 (s, 3H), 1.42–1.58 (m, 3H), 0.88 (d, 3H, $J = 6.6$ Hz), 0.83 (d, 2H, $J = 6.6$ Hz); ¹³C NMR (DMSO-d₆), δ ppm: 174.4, 171.6, 169.7, 158.2, 130.6, 130.3, 113.9, 55.4, 54.7, 51.3, 41.4, 36.9, 24.6, 23.5, 22.9, 22.1; HRMS (ESI⁺): calcd: 350.2080 for $\text{C}_{18}\text{H}_{28}\text{N}_3\text{O}_4$; found: 350.2079.

Ac-D-Tyr(Me)-L-Leu-NH₂. ¹H NMR (300 MHz, DMSO-d₆), δ ppm: 8.23 (d, 1H, $J = 6.6$ Hz), 8.16 (d, 1H, $J = 8.4$ Hz), 7.28 (s, 1H), 7.13 (d, 2H, $J = 8.1$ Hz), 6.99 (s, 1H), 6.80 (d, 2H, $J = 8.4$ Hz), 4.37 (q, 1H, $J = 7.2$ Hz), 4.05 (m, 1H), 3.70 (s, 3H), 2.75 (m, 2H), 1.80 (s, 3H), 1.34 (m, 2H), 1.09 (m, 1H), 1.03 (d, 3H, $J = 6.3$ Hz), 0.83 (d, 2H, $J = 6.3$ Hz); ¹³C NMR (DMSO-d₆), δ ppm: 174.7, 171.7, 170.1, 158.3, 130.6, 129.6, 113.9, 55.6, 55.3, 51.1, 37.1, 24.2, 23.6, 22.8, 21.6; HRMS (ESI⁺): calcd: 350.2080 for $\text{C}_{18}\text{H}_{28}\text{N}_3\text{O}_4$; found: 350.2078.†

Preparation of vesicles/supramolecular aggregates. A poly-disperse suspension of supramolecular aggregates (typically bilayer vesicles, given the pH of the sample) was obtained by simply dissolving the amphiphile/surfactant in aqueous solution, gently stirring and sonicating for 10 min. A concentrated stock solution of the amphiphile/surfactant was prepared in water and adjusted to the pH of the experiment with a 1 M solution of NaOH. In parallel, a concentrated stock solution of the amino acid reactant was prepared in a buffer at the desired pH. The type of buffer used depended on the experimental pH: 2-(*N*-morpholino)ethanesulphonic acid (MES) buffer was used for acidic pH (6.5); 3-(*N*-morpholino)propane sulphonic acid (MOPS) buffer was used for intermediate pHs (7.2–8) and PIPPS buffer for basic pH (9). These two stock solutions were then mixed in convenient proportions to obtain the different concentrations analyzed. The whole process was carried out at the melting temperature (T_m) of the amphiphile/surfactant.

CVC determination. CVC values for decanoic acid were measured using a published protocol.^{16,18,39} A serial dilution of vesicle samples was prepared following the general protocol, starting from a concentrated amphiphile solution of decanoic acid. Merocyanine 570 (hydrophobic dye, 10 μM) was added to every sample after the vesicles were formed and just before spectrophotometric analysis was carried out (using a UV-Vis spectrophotometer Perkin Elmer Lambda 20). Wavelength scanning was performed between 350 and 650 nm, and the CVC was determined as the concentration of amphiphile at which the absorbance ratio between the 570 nm and 530 nm wavelengths increased above the baseline (~ 0.2).

Peptide-forming reactions

Experiments in aqueous media with fatty acids. After the formation of vesicles in the presence of a nucleophilic reagent (5 mM H-L-Leu-NH₂, H-L-Ala-NH₂, H-Gly-NH₂, or H-L-Leu-OH), a 20 mM stock solution of the oxazolone (50 μL) in acetonitrile was added to every sample to give a 1 mM solution and the medium was allowed to react for 2 h at room temperature. Fatty acids were then removed by extraction from the reaction medium with hexane before HPLC analysis. To this aim, samples were acidified with formic acid and then the surfactant was extracted twice with hexane (1 mL). The mixture was gently stirred with a vortex and the upper layer was removed with a pipette. Residual hexane was removed under reduced pressure before analysis by HPLC (Method A).

Experiments in aqueous media with 1,2-dioleoyl-*sn*-glycerol-3-phosphate (DOPA). DOPA containing reaction media were prepared from a 20 mM stock solution of DOPA to yield a 5 mM H-L-Leu-NH₂ solution in a 100 mM MOPS buffer at pH 8. After reaction of the oxazolone (from a 20 mM stock solution in acetonitrile), attempts to remove the surfactant were made by adding tributylammonium bromide and HCl to acidify the samples. Extraction was done following the general protocol, though an extended period was needed for decantation.

Experiments in aqueous media with sodium dodecyl sulfate (SDS). A 100 mM stock solution of SDS in water was mixed with a stock solution of 10 mM H-Leu-NH₂ in 200 mM PIPPS buffer at pH 9. HPLC analysis was performed after dilution of the samples to an SDS concentration below the CMC (10 mM). A similar protocol was used in the case of dodecyl trimethyl ammonium bromide (DoTAB).

Experiments in organic solvents ($\text{CH}_2\text{Cl}_2/\text{CH}_3\text{CN}$). CH_2Cl_2 and CH_3CN were used in the case of acetic acid (AcA) and decanoic acid (DA), and only CH_3CN in the case of *N*-acetyl-D,L-leucine (Ac-D,L-Leu-OH) and *N*-acetyl-L-leucine (Ac-L-Leu-OH) due to the precipitation of solids in CH_2Cl_2 . After the reaction, 6 mM HCl (1 mL) was added to 1 mL samples withdrawn from the reaction medium, as well as 10 mL of hexane. The mixture was then gently stirred with a vortex and the upper layer was removed with a pipette. Hexane addition and extraction was repeated. Finally, residual hexane was removed by evaporation under reduced pressure.

Experiment with Ac-Leu-oxazolone 2b. A 1 mM Ac-L-Tyr(Me)-OH and 5 mM H-L-Leu-NH₂ solution was prepared in CH_3CN (5 mL). The reaction was initiated by the addition of Ac-Leu oxazolone (2 mM final concentration). 1 mL samples of the reaction medium were withdrawn, acidified with 6 mM HCl (1 mL) and extracted once with hexane before measuring the products of any possible reaction of Ac-Tyr(Me)-OH by HPLC (Method A) and detection at 273 nm. This procedure was repeated after one hour, three days and one week (Fig. S4†) allowing the identification of unchanged Ac-Tyr(Me)-OH (retention time 8.7 min) and Ac-L-Tyr(Me)-L-Leu-NH₂ (retention time 14.3 min) as a single L,L stereoisomer. The solution was also analyzed using HPLC-MS confirming the identification of the products as the two peaks corresponding to the isomers of Ac-Leu-Leu-NH₂ (m/z 308.20 [$\text{M} + \text{Na}^+$], retention times 1.91 and



2.01 min; Method B) and that of Ac-Tyr(Me)-Leu-NH₂ (*m/z* 372.19 [M + Na⁺], retention time 1.99 min).

Partition of dipeptide between solution and surfactant

The Langmuir–Blodgett monolayer technique (Kibron Micro-Trough XS) was used to elucidate the interactions between the Ac-L-Tyr(Me)-L-Leu-NH₂ dipeptide and the lipid. A monolayer of fatty acid was spread on a well with 1 mL of buffer at the corresponding pH registering an initial stabilized pressure (Π_0). 10 μ L of a 5 mM Ac-L-Tyr(Me)-L-Leu-NH₂ dipeptide stock solution in MeOH was then added from the water face. After MeOH evaporation, any surface pressure increase resulted from a migration of the hydrophobic solute to the monolayer, stabilizing the signal at a new pressure (Π_1). The difference between these two surface pressures ($\Delta\Pi$) was recorded and compared to the Π_0 in every individual experiment.

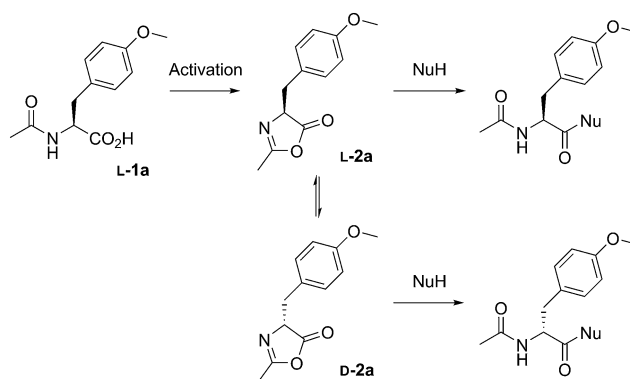
Binding of Ac-L-Tyr(Me)-L-Leu-NH₂ to the vesicles

This experimental assay is based on the quenching action of acrylamide, which decreases the intrinsic fluorescence of Tyr residues as a function of its local concentration. Acrylamide was taken from a 6.25 M stock solution and added in 4 μ L portions (5 additions, from 0 mM of acrylamide to 250 mM in the final volume) to the control and 1 mL samples containing vesicles (which were previously prepared, following the general procedure, in the presence of 1 μ M Ac-L-Tyr(Me)-L-Leu-NH₂ dipeptide). Fluorescence was monitored at a single wavelength (ex: 273 nm and em: 340 nm) after every addition. In parallel, a dilution control was carried out, adding the same volume of aqueous buffer instead of acrylamide solution. The fluorescence obtained as a result of the addition of only buffer (F_0) was divided by the fluorescence obtained as a result of the addition of acrylamide (F) and compared against the concentration values of acrylamide added.

Results

Experiments in aqueous media

We extensively studied the outcome of reactions taking place when a racemic mixture of hydrophobic 5(4*H*)-oxazolone enantiomers (L-2a and D-2a), derived from Ac-L-Tyr(Me)-OH L-1a (Scheme 1), was put in contact with H-L-Leu-NH₂ in buffered media. Owing to its chiral instability, the 5(4*H*)-oxazolones undergo a kinetically stereoselective reaction with the nucleophiles, which have a fixed configuration,³⁶ meaning that the timescale of the epimerization process is similar to or faster than the subsequent reaction and that the configuration of products is kinetically controlled. Reactions producing only peptide diastereomers and racemic Ac-Tyr(Me)-OH as a result of hydrolysis were carried out under different pH conditions and in the presence of different amounts of amphiphiles and surfactants. For each case, both the total dipeptide yield (relative to the amount of starting material) and the ratio of diastereomers were determined. Fig. 1 shows the characteristic results obtained when carboxylic acids are employed (in this case: decanoic acid (DA) at pH 7.2, but see Fig. 2 and Table S1 in



Scheme 1 The reactions performed in this work from racemic mixtures of the 5(4*H*)-oxazolone enantiomers (L-2a and D-2a) with nucleophiles (NuH = H-L-Leu-NH₂, H-L-Ala-NH₂, H-Gly-NH₂, or H-L-Leu-OH) in aqueous media, leading to the corresponding dipeptides as mixtures of enantiomers (Ac-L-Tyr(Me)-Gly-NH₂ and Ac-D-Tyr(Me)-Gly-NH₂) or diastereomers (Ac-L-Tyr(Me)-L-Leu-NH₂ and Ac-D-Tyr(Me)-L-Leu-NH₂, Ac-L-Tyr(Me)-L-Ala-NH₂ and Ac-D-Tyr(Me)-L-Ala-NH₂, or Ac-L-Tyr(Me)-L-Leu-OH and Ac-D-Tyr(Me)-L-Leu-OH). Although prepared from chiral Ac-L-Tyr(Me)-OH (L-1a), the oxazolone was present in a racemic state due to its chiral instability.³⁶

the ESI[†] for further data). Typically, there is a 2 to 3-fold increase in the overall yield of dipeptide, together with a reversion in its diastereomeric ratio (from a preference for homo-chiral forms, in solution,³⁶ toward more heterochiral ones).

As is apparent from Fig. 1, this double effect only happens provided that the concentration of fatty acid is above its critical aggregation concentration (CAC) value (the critical vesicular concentration, CVC, in this case), which was measured independently (Fig. S2 in the ESI[†]), and reaches a plateau soon after that critical value is crossed. Those two independent observations suggest that the reactions in which hydrophobic substrates are involved take place in a different environment, whenever fatty acid aggregates offer this possibility. The double effect may well depend on the fact that the nucleophilic reaction

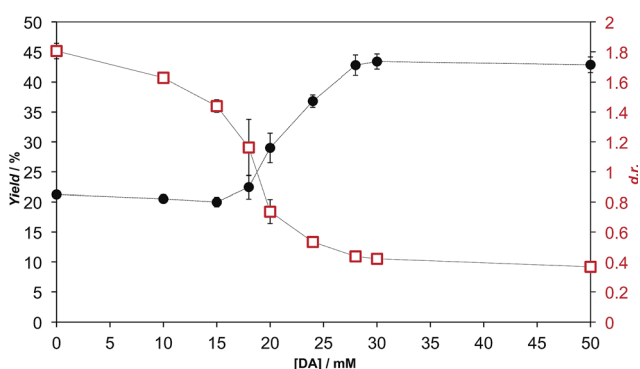


Fig. 1 Formation of peptide diastereomers Ac-L-Tyr(Me)-L-Leu-NH₂ and Ac-D-Tyr(Me)-L-Leu-NH₂ by the reaction of 1 mM racemic 5(4*H*)-oxazolone 2a with 5 mM H-L-Leu-NH₂ in 100 mM MOPS buffer (pH 7.2), at room temperature, in the presence of different concentrations of decanoic acid (DA). The effects are reported in terms of the relative total yield of the dipeptide (left axis, black filled circles) and of the diastereomeric ratio (right axis, red open squares).



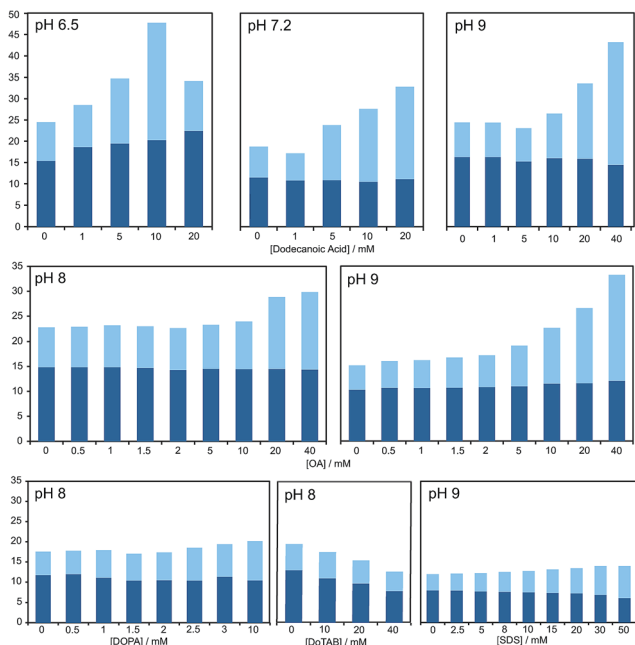


Fig. 2 Bar graph representing the total yield of dipeptide (full bar size) and stereoselectivity (chiral conversion %: Ac-L-Tyr(Me)-L-Leu-NH₂, in dark blue; and Ac-L-Tyr(Me)-L-Leu-NH₂, in light blue) which are obtained in reactions carried out at room temperature, in aqueous buffer (100 mM) at various pH values, starting from Ac-Tyr(Me) oxazolone (1 mM), in the presence of H-L-Leu-NH₂ (5 mM) and diverse amphiphile/surfactant concentrations. Each bar corresponds to a single experiment; error fluctuation was estimated with the control samples, without amphiphile (based on d.r. data and calculated for each pH value – see Table S1†) and remains within the range of 12–18% for all cases. OA stands for oleic acid. For a more exhaustive report of the results, see main text and Table S1 in the ESI.†

partners have a strong enough affinity for the hydrophobic environment (see data for alaninamide, glycinamide and leucine, instead of leucinamide, at the end of Table S1 in the ESI†). It is difficult to explain, otherwise, how the minimal variations in the chemical nature of the reactants (the reduced aliphatic side-chain in the first two cases, or the higher polarity in the third, zwitterionic one) make such a difference.

Additional assays were carried out to complement the previous results and check whether the product of the reaction co-localizes with the hydrophobic supramolecular domains. More specifically, a series of quenching experiments with acrylamide (Fig. S3 in ESI†) show that when DA vesicles are present in the medium, the fluorescent signal coming from the Tyr ring of the dipeptide could not be screened down, as occurs in free solution. Furthermore, data collected from Langmuir trough experiments (Fig. S4 in the ESI†) also show that the addition of the dipeptide to DA monolayers with different initial surface pressures (π_0) induced an increase in the surface pressure ($\Delta\pi$), which results from the adsorption of the dipeptide to the monolayer.

Our HPLC results were consistent over a relatively wide range of concentrations, carboxylic acid species and pH values (see Fig. 2 and Table S1 in the ESI†). Nevertheless, stronger

effects were typically obtained at pH values that do not largely exceed the pK_a of the corresponding fatty acid in the charged aggregate environment, which suggests that the vesicles had the optimal supra-molecular configuration to achieve a more significant influence on the reaction (at lower pH less reliable results were collected, owing probably to the formation of droplets).

A series of additional control experiments was performed to check the specificity of our results with regard to the type of amphiphilic/surfactant molecule employed. Since fatty acid vesicles are, overall, negatively charged (due to the partial deprotonation of their monomeric units), we tried with negatively charged species containing aliphatic chains of a similar length, like the phosphatidic acid DOPA (1,2-dioleoyl-*sn*-glycero-3-phosphoric acid) or sodium dodecyl sulfate (SDS).

As reported in Fig. 2 and Table S1 (ESI†), no detectable effects on the yield of the reaction were found in those cases, even if we did observe changes in the diastereomeric ratios, which were altogether absent if a positively charged surfactant (like dodecyltrimethylammonium bromide, DoTAB) was used. Although an explanation for this inverted trend in the chiral nature of the products (preference for *D*-forms in the majority of cases when supramolecular structures are present) is not within the scope of this study, it is nevertheless useful as an indicator of the environment in which the reaction actually takes place. This view is also supported by the known effect of low polarity solvents favouring heterochiral adducts of 5(*4H*)-oxazolones.⁴⁰ Finally, it should be mentioned that mixtures of decanoic acid with decanol (previously reported to confer higher stability to the resulting vesicles in aqueous solution)¹⁴ were tried, too, but the effects on the reaction were clearly diminished.

Experiments in organic solvents

All the results obtained in aqueous buffered solution indicate that the pathway through which fatty acids play a role does not simply correspond to the usual activity of hydrophobic assemblies on chemical reactivity. Indeed, a specific behaviour of carboxylic acids at pH values inducing the formation of bilayers might be related to the activity of their polar head groups, combined in both the neutral and anionic forms, at the lipid-water interface of the supramolecular structure (Fig. S1 in the ESI†). In order to determine what kind of chemical activity is effectively responsible for this behaviour, our reaction was also studied in organic solvents, in the presence of decanoic acid and acetic acid. Both acids induced a substantial increase in reaction rate in two different solvents, CH₂Cl₂ (Fig. 3) and CH₃CN, and the hydrophobic chain of decanoic acid had no significant additional influence (see Table 1). The more acidic Ac-Leu-OH was studied in CH₃CN to avoid solubility issues and had a slightly improved activity at low concentrations (2 mM), compared to acetic acid and decanoic acid under the same conditions (Table S2 in ESI†). Therefore, the activity of fatty acids does not depend on the presence of a hydrophobic aliphatic chain, which indicates that it is the result of the participation of the carboxylic acid moiety.



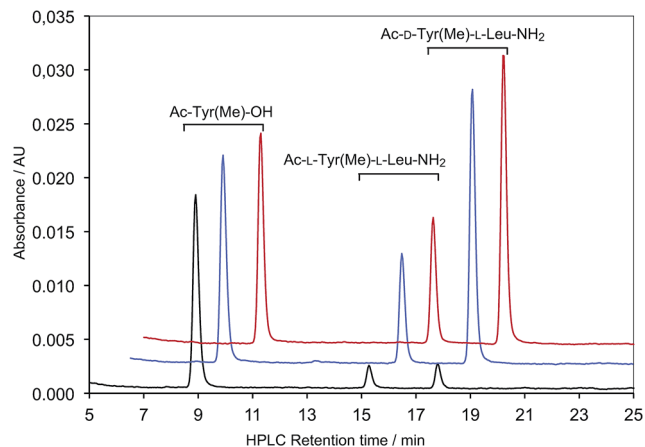


Fig. 3 HPLC monitoring of the reaction (samples withdrawn after 1 h) of Ac-L-Tyr(Me)-5(4*H*)-oxazolone **2a** (ca. 1 mM) with H-L-Leu-NH₂ (5 mM) carried out in CH₂Cl₂ (control, black) or in the presence of added carboxylic acids (2 mM): decanoic acid (red) or acetic acid (blue).

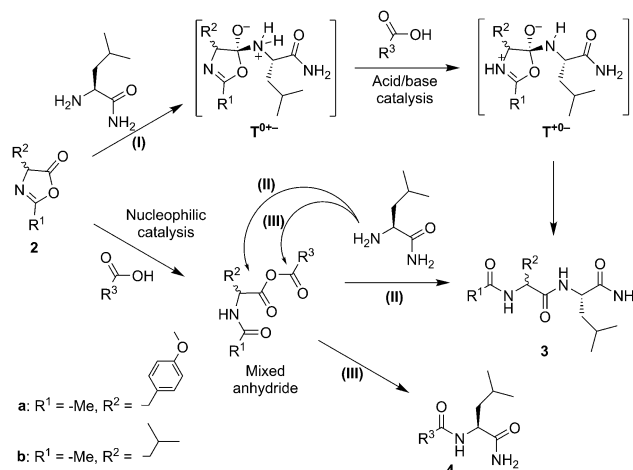
Table 1 Yield and stereoselectivity (shown as diastereomeric ratio, d.r.) in selected reactions carried out in organic solvent for Ac-Tyr(Me) oxazolone (1 mM) in the presence of 5 mM H-Leu-NH₂ and carboxylic acid reagents at room temperature for 1 h. For an exhaustive description of the results see Table S2 of the ESI

Solvent	Carboxylic acid	Yield/%	d.r.
CH ₂ Cl ₂	Control without acid	17.4	0.84
CH ₂ Cl ₂	10 mM acetic acid	86.2	0.49
CH ₂ Cl ₂	10 mM DA	86.9	0.54
CH ₃ CN	Control without acid	2.7	0.62
CH ₃ CN	10 mM acetic acid	45.3	0.44
CH ₃ CN	10 mM DA	55.7	0.46

Discussion

Our experimental survey proves that fatty acid aggregates increase the yield of peptide formation from a hydrophobic 5(4*H*)-oxazolone and leucinamide. The product of the reaction was also demonstrated to have an affinity for the lipid phase. The favourable effect on the yield is specific of surfactants that bear a carboxylic acid moiety. This effect is accompanied by a reversal of stereoselectivity observed only above the CAC, which requires a more elaborate explanation but is beyond the scope of the present work.

In organic solvents a similar preference for heterochiral products was observed, together with a rate enhancement promoted by carboxylic acids. In these media, carboxylic acids can either act as nucleophilic catalysts or acid–base catalysts (Scheme 2). Clues attesting to acid–base catalysis can neither be expected from the accumulation of intermediates nor from the detection of side-products. However, thanks to the presence of two electrophilic sites in the mixed anhydride intermediate, a transfer of activation from the 5(4*H*)-oxazolone **2a** to the carboxylic acid R³-COOH leading to amide side-product **4** (rather than to the dipeptide Ac-Tyr(Me)-Leu-NH₂ **3**) can, in contrast, be predicted during the nucleophilic pathway. This



Scheme 2 Potential chemical pathways of acid–base (I) and nucleophilic (II and III) catalysis (during the aminolysis of 5(4*H*)-oxazolones **2a** and **2b** by leucinamide in organic solvents) suggested as a way to account for the activity of carboxylic acids R³COOH (which include acetic acid, decanoic acid and the acylated α -amino acid Ac-D,L-Leu-OH and optically active Ac-L-Leu-OH and Ac-L-Tyr(Me)-OH).

would come as a result of path (III), in addition to the aminolysis yielding product **3**, as previously obtained from path (II). To assess the extent of nucleophilic catalysis, an experiment was carried out starting from 2 mM leucine 5(4*H*)-oxazolone **2b** in acetonitrile and using 1 mM Ac-L-Tyr(Me)-OH **1a** as a carboxylic acid component (R³COOH). The formation of Ac-L-Tyr(Me)-L-Leu-NH₂ dipeptide was observed to be limited to 1% (with respect to **1a**) after 1 h, and slowly increased up to 5% after 3 days (Fig. S5 in the ESI†). Importantly, Ac-L-Tyr(Me)-L-Leu-NH₂ dipeptide was formed as a single stereoisomer, ruling out any reaction path occurring through the 5(4*H*)-oxazolone **2a**, and supporting the mixed anhydride one (III). This experiment, in fact, suggests that the mixed anhydride pathway is a slowly proceeding process.

The fast and highly efficient increase in the rate (ca. 10-fold) observed by adding 2 mM Ac-Leu-OH (as R³COOH) to 1 mM Ac-Tyr(Me) 5(4*H*)-oxazolone **2a** must therefore be considered to be the result of the acid–base-catalysed path (I). The faster reaction of 5(4*H*)-oxazolone **2a**, compared to that of the mixed anhydride, is not surprising if one takes into account two key points: (i) the lesser importance of steric hindrance for the nucleophilic attack at the 5(4*H*)-oxazolone carbonyl group, relative to that occurring at the two reacting centres of the mixed anhydride; and (ii) the better leaving group ability of the amide oxygen (p*K*_a ~ -1.0 in water for the conjugate acid) relative to that of a carboxylate group (p*K*_a ~ 4–5 in water for carboxylic acids – although a significantly higher value is expected in the anionic environment of the aggregate).

The occurrence of acid–base catalysis of any kind is directly connected to the need for proton transfer during the course of the reaction, given the large change in p*K*_a of the reacting groups involved.^{41,42} In path (I) two proton transfers are needed for conversion into the product. The tetrahedral intermediate T⁰⁺ is produced by the direct addition of the nucleophile to the



5(4*H*)-oxazolone. The need for reaching T^{+0-} by proton transfer before any conversion into products takes place can account for the catalysis by carboxylic acid observed in organic solvents (Scheme 2, path (I)). The most straightforward process is a pathway starting from this intermediate by protonation of the nitrogen atom of the leaving group by the carboxylic acid, thus facilitating breakdown. The alternative possibility of the carboxylate anion acting as a base can be ruled out by the fact that the base, H-Leu-NH₂, present at the same time in the medium (in 5 mM concentration) was not sufficient to promote the reaction (Fig. 3, control). An additional advantage of carboxylic acids could be their ability to act as both an acid and base in a bifunctional way,⁴³ which could make the decomposition of the tetrahedral intermediate easier by the direct conversion of T^{0+-} into T^{+0-} .

The results reported here for the reactions carried out in aqueous media could be explained by the possibility that fatty acid supramolecular aggregates have a catalytic effect similar to the micellar catalysis already demonstrated for bimolecular reactions between hydrophobic substrates (*i.e.*, selective concentration effect in a reduced volume).^{44,45} The need for a concentration of surfactant exceeding the CAC (in order to have an observable effect) indeed supports the involvement of hydrophobic aggregates. However, this explanation is not sufficient to account for the highly specific behaviour that fatty acid aggregates (as compared to other lipid or surfactant aggregates) were observed to have, inducing significant increases in the yield of peptide formation reactions from 5(4*H*)-oxazolones. A more comprehensive explanation of all the data collected in this work, coherent with the results obtained in organic solvents, would include an acid–base type of catalysis taking place at the lipid–water interface of the supramolecular structures (Scheme 3), where the polar heads of the carboxylic acids undergo continuous protonation–deprotonation dynamics. Precisely the same ‘chemical trick’ that enables the stability of fatty acid vesicles as prebiotic compartments or microreactors (a balanced combination of neutral and anionic forms of the monomer) could be the critical feature to account for their catalytic power.

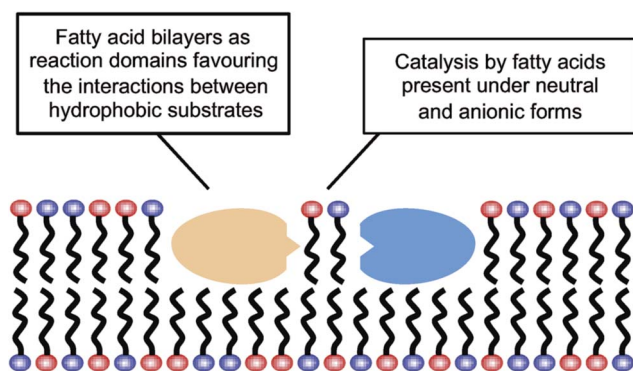
In this context, it is noteworthy that most aminolysis processes required for peptide syntheses involve large changes

in the pK_a of the reacting centres and, therefore, are susceptible to acid–base catalysis in a way similar to the reactions of 5(4*H*)-oxazolones. Bilayers made of fatty acids demonstrate that unique ability at neutral pH, which is not realised for other types of lipid compartments, like the phospholipid-based boundaries of modern cells. So fatty acids, in addition to providing a different local environment for the reaction (through their spontaneous assembly into supramolecular structures), could also act as acid–base catalysts for the formation of hydrophobic peptides, and potentially for any other chemical processes in which rate-determining proton transfer is required. These peptides, in turn, would likely contribute to membrane stability and would foster growth and potential reproduction of the vesicles. Therefore, one can envision a scenario in which lipid and protein precursors (*i.e.* fatty acids and hydrophobic peptides) would come together and mutually reinforce each other, in an integrated approach to the origin of the first functional protocells. This incipient stage is fully congruent with the modern structure of biological membranes, which consist of phospholipids and also of other hydrophobic components, most prominently membrane-associated or transmembrane proteins. Accordingly, heterogeneous compartments, made of much simpler but already diverse kinds of hydrophobic compounds, would be present from the earliest phases of evolution, preceding the selection of current biomembranes.

More broadly speaking, this scenario is in line with recent perspectives on the problem of how chemistry could open the way towards biological phenomena, within a *systems* framework.^{3,4} The main idea underlying this new conception is that life, from its first steps, cannot be reduced to the chemistry of a single type of compound (*e.g.*, nucleotide chemistry). Instead, prebiotic systems (taken as integrated entities, not just ‘populations of molecules’) ought to encompass several types of biomolecules – or rather, biomolecule *precursors* – in continuous interaction. In other words, the whole process of the origins of life would be understood as the evolutionary development of heterogeneous chemical mixtures with diverse emergent functionalities. Complex biopolymers (like the ones making up extant living cells) would be the final outcome of this long evolutionary process, not the starting point. Our results, from experiments carried out precisely with two main kinds of biomolecular precursor, fatty acids and amino acids, support this alternative view of the problem, in line with other recent experimental reports^{32,34} that highlight the importance of connecting peptide chemistry and compartments.

Acknowledgements

SMS and KRM received support from the Basque Government (IT 590-13 Grant) and from the Spanish Ministry of Economía y Competitividad (FFI2014-52173-P Grant). DB and RP thank the Agence Nationale de la Recherche for supporting the PeptiSystems project (grant ANR-14-CE33-0020). The authors also thank COST Action CM1304 (Emergence and Evolution of Complex Chemical Systems) for eliciting the collaboration between our groups and for supporting this project through a Short-Term



Scheme 3 Fatty acids' dual role in peptide bond formation.



Scientific Mission for SMS, who holds a PhD Grant from the University of the Basque Country (UPV/EHU).

Notes and references

‡ HPLC peaks of all the other dipeptide diastereomers used in this work were identified by comparison with authentic samples prepared earlier.³⁶

- R. F. Ludlow and S. Otto, *Chem. Soc. Rev.*, 2008, **37**, 101–108.
- G. von Kiedrowski, S. Otto and P. J. Herdewijn, *J. Syst. Chem.*, 2010, **1**, 1–6.
- K. Ruiz-Mirazo, C. Briones and A. de la Escosura, *Chem. Rev.*, 2014, **114**, 285–366.
- B. H. Patel, C. Percivalle, D. J. Ritson, C. D. Duffy and J. D. Sutherland, *Nat. Chem.*, 2015, **7**, 301–307.
- S. Pizzarello and E. Shock, *Cold Spring Harbor Perspect. Biol.*, 2010, **2**, a002105.
- H. J. Cleaves, J. H. Chalmers, A. Lazcano, S. L. Miller and J. L. Bada, *Origins Life Evol. Biosphere*, 2008, **38**, 105–115.
- L. P. Dworkin, D. W. Deamer, S. A. Sandford and L. Allamandola, *Proc. Natl. Acad. Sci. U. S. A.*, 2001, **98**, 815–819.
- T. M. McCollom, G. Ritter and B. R. Simoneit, *Origins Life Evol. Biospheres*, 1999, **29**, 153–166.
- G. Danger, R. Plasson and R. Pascal, *Chem. Soc. Rev.*, 2012, **41**, 5416–5429.
- R. Pascal, L. Boiteau and A. Commeyras, *Top. Curr. Chem.*, 2005, **259**, 69–122.
- G. Danger, A. Michaut, M. Bucchi, L. Boiteau, J. Canal, R. Plasson and R. Pascal, *Angew. Chem., Int. Ed.*, 2013, **52**, 611–614.
- W. R. Hargreaves and D. W. Deamer, *Biochemistry*, 1978, **17**, 3759–3768.
- J. M. Gebicki and M. Hicks, *Nature*, 1973, **243**, 232–234.
- C. L. Apel, D. W. Deamer and M. N. Mautner, *Biochim. Biophys. Acta, Biomembr.*, 2002, **1559**, 1–9.
- M. M. Hanczyc and J. W. Szostak, *Curr. Opin. Chem. Biol.*, 2004, **8**, 660–664.
- T. Namani and P. Walde, *Langmuir*, 2005, **21**, 6210–6219.
- K. Morigaki and P. Walde, *Curr. Opin. Colloid Interface Sci.*, 2007, **12**, 75–80.
- S. E. Maurer, D. W. Deamer, J. M. Boncella and P.-A. Monnard, *Astrobiology*, 2009, **9**, 979–987.
- S. E. Maurer and P.-A. Monnard, *Entropy*, 2011, **13**, 466–484.
- I. A. Chen and P. Walde, *Cold Spring Harbor Perspect. Biol.*, 2010, **2**, a002170.
- P. Walde, H. Umakoshi, P. Stano and F. Mavelli, *Chem. Commun.*, 2014, **50**, 10177–10197.
- J. Blain and J. W. Szostak, *Annu. Rev. Biochem.*, 2014, **83**, 615–640.
- H. Yanagawa, Y. Ogawa, K. Kojima and M. Ito, *Origins Life Evol. Biosphere*, 1988, **18**, 179–207.
- D. W. Armstrong, R. Seguin, C. J. McNeal, R. D. Macfarlane and J. H. Fendler, *J. Am. Chem. Soc.*, 1978, **100**, 4605–4606.
- A. Brack, *Origins Life Evol. Biosphere*, 1987, **17**, 367–379.
- Z. Liu, D. Beauflis, J.-C. Rossi and R. Pascal, *Sci. Rep.*, 2014, **4**, 7440.
- M. Blocher, D. Liu, P. Walde and P. L. Luisi, *Macromolecules*, 1999, **32**, 7332–7334.
- M. Blocher, D. Liu and P. L. Luisi, *Macromolecules*, 2000, **33**, 5787–5796.
- M. Blocher, T. Hitz and P. L. Luisi, *Helv. Chim. Acta*, 2001, **84**, 842–848.
- T. Hitz, M. Blocher, P. Walde and P. L. Luisi, *Macromolecules*, 2001, **34**, 2443–2449.
- H. H. Zepik, S. Rajamani, M.-C. Maurel and D. Deamer, *Origins Life Evol. Biosphere*, 2007, **37**, 495–505.
- A. Grochmal, L. Prout, R. Makin-Taylor, R. Prohens and S. Tomas, *J. Am. Chem. Soc.*, 2015, **137**, 12269–12275.
- R. Furuuchi, E. I. Imai, H. Honda, K. Hatori and K. Matsuno, *Origins Life Evol. Biosphere*, 2005, **35**, 333–343.
- K. Adamala and J. W. Szostak, *Nat. Chem.*, 2013, **5**, 495–501.
- Z. R. Todd and C. H. House, *Astrobiology*, 2014, **14**, 859–865.
- D. Beauflis, G. Danger, L. Boiteau, *et al.*, *Chem. Commun.*, 2014, **50**, 3100–3102.
- F. M. F. Chen, K. Kuroda and N. L. Benoiton, *Synthesis*, 1979, 230–232.
- F. M. F. Chen and N. L. Benoiton, *Int. J. Pept. Protein Res.*, 2009, **30**, 683–688.
- J. L. Cape, P.-A. Monnard and J. M. Boncella, *Chem. Sci.*, 2011, **2**, 661–671.
- N. L. Benoiton, K. Kuroda and F. M. F. Chen, *Tetrahedron Lett.*, 1981, **22**, 3361–3364.
- W. P. Jencks, *Acc. Chem. Res.*, 1976, **9**, 425–432.
- W. P. Jencks, *Acc. Chem. Res.*, 1980, **13**, 161–169.
- A. C. Satterthwait and W. P. Jencks, *J. Am. Chem. Soc.*, 1974, **96**, 7031–7044.
- C. A. Bunton, F. Nome, F. H. Quina and L. S. Romsted, *Acc. Chem. Res.*, 1991, **24**, 357–364.
- S. Otto and J. B. F. N. Engberts, *Org. Biomol. Chem.*, 2003, **1**, 2809–2820.

